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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2008; month=4; day=16; hr=16; min=11; sec=13; ms=74;]

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Application No: 10500671

Version No: 3.0

Input Set:**Output Set:****Started:** 2008-04-01 15:52:20.620**Finished:** 2008-04-01 15:52:22.048**Elapsed:** 0 hr(s) 0 min(s) 1 sec(s) 428 ms**Total Warnings:** 18**Total Errors:** 0**No. of SeqIDs Defined:** 21**Actual SeqID Count:** 21

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| W 213 | Artificial or Unknown found in <213> in SEQ ID (18) |
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SEQUENCE LISTING

<110> Sera, Takashi

<120> Nuclear-Envelope and Nuclear-Lamina Binding Chimeras for
Modulating Gene Expression

<130> 109845-163

<140> 10500671

<141> 2008-04-01

<160> 21

<170> PatentIn version 3.3

<210> 1

<211> 25

<212> PRT

<213> Artificial

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<223> Zinc finger domain

<220>

<221> MISC_FEATURE

<222> (2)..(5)

<223> Amino acids 2-5 are Xaa wherein Xaa = any amino acid, and up to
two amino acids can be missing.

<220>

<221> MISC_FEATURE

<222> (7)..(18)

<223> Xaa can be any amino acid

<220>

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<222> (20)..(24)

<223> Amino acids 20-24 are Xaa wherein Xaa = any amino acid, and up to
two amino acids can be missing.

<400> 1

Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

1 5 10 15

Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His

20 25

<210> 2

<211> 32

<212> PRT

<213> Artificial

<220>
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 <223> Xaa can be any amino acid

<220>
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 <222> (5)..(8)
 <223> Amino acids 5-8 are Xaa wherein Xaa = any amino acid, and up to two amino acids can be missing

<220>
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 <222> (10)..(14)
 <223> Xaa can be any amino acid

<220>
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 <222> (15)..(15)
 <223> Amino acid 15 is Z(-1) wherein Z(-1) = Arg, Lys, Gln, Asn, Thr, Met, Leu, Ile, Glu or Asp.

<220>
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 <222> (16)..(16)
 <223> Xaa can be any amino acid

<220>
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 <222> (17)..(17)
 <223> Amino acid 17 is Z2 wherein Z2 = Ser, Arg, Asn, Gln, Thr, Val, Ala, Asp or Glu.

<220>
 <221> MISC_FEATURE
 <222> (18)..(18)
 <223> Amino acid 18 is Z3 wherein Z3 = His, Lys, Asn, Gln, Ser, Ala, Val, Thr, Asp, or Glu

<220>
 <221> MISC_FEATURE
 <222> (19)..(20)
 <223> Xaa can be any amino acid

<220>
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 <222> (21)..(21)
 <223> Amino acid 21 is Z6 wherein Z6 = Arg, Lys, Gln, Asn, Thr, Tyr, Leu, Ile, Met, Glu or Asp.

<220>
 <221> MISC_FEATURE
 <222> (23)..(27)

<223> Amino acids 23-27 are Xaa wherein Xaa = any amino acid, adn up to two amino acids can be missing.

<220>

<221> MISC_FEATURE

<222> (29)..(32)

<223> Xaa can be any amino acid

<400> 2

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa
20 25 30

<210> 3

<211> 28

<212> PRT

<213> Artificial

<220>

<223> Zinc finger domain

<220>

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<222> (13)..(13)

<223> Amino acid 13 is Z(-1) wherein Z(-1) = Arg, Lys, Gln, Asn, Thr, Met, Leu, Ile, Glu or Asp.

<220>

<221> MISC_FEATURE

<222> (15)..(15)

<223> Amino acid 15 is Z2 wherein Z2 = Ser, Arg, Asn, Gln, Thr, Val, Ala, Asp or Glu.

<220>

<221> MISC_FEATURE

<222> (16)..(16)

<223> Amino acid 16 is Z3 wherein Z3 = His, Lys, Asn, Gln, Ser, Ala, Val, Thr, Asp or Glu.

<220>

<221> MISC_FEATURE

<222> (19)..(19)

<223> Amino acid 19 is Z6 wherein Z6 = Arg, Lys, Gln, Asn, Thr, Tyr, Leu, Ile, Met, Glu or Asp.

<400> 3

Pro Tyr Lys Cys Pro Glu Cys Gly Lys Ser Phe Ser Xaa Ser Xaa Xaa
1 5 10 15

Leu Gln Xaa His Gln Arg Thr His Thr Gly Glu Lys
20 25

<210> 4

<211> 5

<212> PRT

<213> Artificial

<220>

<223> Synthetic flexible linker peptide for linking together
multi-finger zinc finger domains

<400> 4

Gly Gly Gly Gly Ser
1 5

<210> 5

<211> 11

<212> PRT

<213> Human immunodeficiency virus

<220>

<221> MISC_FEATURE

<222> (1)..(11)

<223> HIV Tat protein domain

<400> 5

Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
1 5 10

<210> 6

<211> 9

<212> DNA

<213> Human immunodeficiency virus

<220>

<221> misc_feature

<222> (1)..(9)

<223> HIV DNA Binding Domain

<400> 6

gcagaagcc

9

<210> 7

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<212> DNA

<213> Artificial

<220>

<223> DNA target sequence

<400> 7

gtgtgggtga gtgagtgtg

19

<210> 8

<211> 19

<212> DNA

<213> Artificial

<220>

<223> DNA target sequence

<400> 8

ggggctgggg gcggtgtct

19

<210> 9

<211> 7

<212> PRT

<213> Simian virus 40

<220>

<221> MISC_FEATURE

<222> (1)..(7)

<223> Peptide from SV40 large T antigen

<400> 9

Pro Lys Lys Lys Arg Lys Val

1 5

<210> 10

<211> 16

<212> PRT

<213> Artificial

<220>

<223> Peptide, residues 43-58 of the Antennapeida homeodomain protein

<220>

<221> MISC_FEATURE

<222> (1)..(16)

<223> Peptide, residues 43-58 of the Antennapeida homeodomain protein

<400> 10

Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys

1 5 10 15

<210> 11

<211> 34

<212> PRT
<213> Herpes Simplex Virus

<220>
<221> MISC_FEATURE
<222> (1)..(34)
<223> Residues 267-300 of the HSV VP22 protein

<400> 11

Asp Ala Ala Thr Ala Thr Arg Gly Arg Ser Ala Ala Ser Arg Pro Thr
1 5 10 15

Glu Arg Pro Arg Ala Pro Ala Arg Ser Ala Ser Arg Pro Arg Arg Pro
20 25 30

Val Glu

<210> 12
<211> 11
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide modeled after the protein transduction domain
of the human immunodeficiency virus TAT protein having cellular
uptake activity

<400> 12

Tyr Ala Arg Ala Ala Ala Arg Gln Ala Arg Ala
1 5 10

<210> 13
<211> 9
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide modeled after the protein transduction domain
of the human immunodeficiency virus TAT protein having cellular
uptake activity, referred to as "R9"

<400> 13

Arg Arg Arg Arg Arg Arg Arg Arg
1 5

<210> 14
<211> 16

<212> PRT

<213> Artificial

<220>

<223> D-penetratin peptide

<220>

<221> MISC_FEATURE

<222> (1)..(16)

<223> All amino acids are in the D-form

<400> 14

Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
1 5 10 15

<210> 15

<211> 16

<212> PRT

<213> Artificial

<220>

<223> Peptide Syn B1 from Antennapedia homeodomain protein

<400> 15

Arg Gly Gly Arg Leu Ser Tyr Ser Arg Arg Arg Phe Ser Thr Ser Thr
1 5 10 15

<210> 16

<211> 10

<212> PRT

<213> Artificial

<220>

<223> L-SynB3 peptide from Antennapedia homeodomain protein

<400> 16

Arg Arg Leu Ser Tyr Ser Arg Arg Arg Phe
1 5 10

<210> 17

<211> 10

<212> PRT

<213> Artificial

<220>

<223> D-SynB3 peptide from Antennapedia homeodomain pro

<220>

<221> MISC_FEATURE

<222> (1)..(10)
<223> All amino acids are in the D-form

<400> 17

Arg Arg Leu Ser Tyr Ser Arg Arg Arg Phe
1 5 10

<210> 18
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<212> PRT
<213> Artificial

<220>
<223> Flag Epitope Peptide

<400> 18

Asp Tyr Lys Asp Asp Asp Asp Lys
1 5

<210> 19
<211> 5
<212> PRT
<213> Artificial

<220>
<223> Artificial peptide linker

<400> 19

Gly Gly Gly Gly Ser
1 5

<210> 20
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<223> Artificial peptide linker

<400> 20

Gly Gly Gly Ser
1

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<223> Artificial peptide linker

<400> 21

Gly Gly Ser

1